OIPE

RAW SEQUENCE LISTING   DATE: 06/04/2002
PATENT APPLICATION: US/09/772,116   TIME: 12:41:11

Input Set : A:\Seqlist.txt
Output Set: N:\CRF3\06042002\I772116.raw

ENTERED

```
 3 <110> APPLICANT: Benjamin, Howard
 4        Findeis, Mark A.
 5        Gefter, Malcolm L.
 6        Musso, Gary
 7        Signer, Ethan R.
 9 <120> TITLE OF INVENTION: METHODS FOR IDENTIFYING COMPOUNDS THAT BIND TO A TARGET
11 <130> FILE REFERENCE: PPI-010CN
13 <140> CURRENT APPLICATION NUMBER: US 09/772116
14 <141> CURRENT FILING DATE: 2001-01-26
16 <150> PRIOR APPLICATION NUMBER: US 08/573786
17 <151> PRIOR FILING DATE: 1995-12-18
19 <160> NUMBER OF SEQ ID NOS: 6
21 <170> SOFTWARE: FastSEQ for Windows Version 4.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 16
25 <212> TYPE: PRT
26 <213> ORGANISM: Artificial Sequence
28 <220> FEATURE:
29 <223> OTHER INFORMATION: motif
31 <220> FEATURE:
32 <221> NAME/KEY: VARIANT
33 <222> LOCATION: 1, 6, 11, 16
34 <223> OTHER INFORMATION: Xaa = any amino acid
36 <220> FEATURE:
37 <221> NAME/KEY: VARIANT
38 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15
39 <223> OTHER INFORMATION: Xaa = Ala or Gly
41 <400> SEQUENCE: 1
42 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
43 1               5                   10                  15
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47 <211> LENGTH: 16
48 <212> TYPE: PRT
49 <213> ORGANISM: Artificial Sequence
51 <220> FEATURE:
52 <223> OTHER INFORMATION: motif
54 <220> FEATURE:
55 <221> NAME/KEY: VARIANT
56 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15
57 <223> OTHER INFORMATION: Xaa = Ala or Gly
59 <400> SEQUENCE: 2
60 Ser Xaa Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Met
61 1               5                   10                  15
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 64 <210> SEQ ID NO: 3
 65 <211> LENGTH: 16
 66 <212> TYPE: PRT
 67 <213> ORGANISM: Artificial Sequence
 69 <220> FEATURE:
 70 <223> OTHER INFORMATION: motif
 72 <220> FEATURE:
 73 <221> NAME/KEY: VARIANT
 74 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15
 75 <223> OTHER INFORMATION: Xaa = Ala or Gly
 77 <400> SEQUENCE: 3
 78 Ser Xaa Xaa Xaa Xaa Lys Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Gln
 79  1               5                   10                  15
 82 <210> SEQ ID NO: 4
 83 <211> LENGTH: 16
 84 <212> TYPE: PRT
 85 <213> ORGANISM: Artificial Sequence
 87 <220> FEATURE:
 88 <223> OTHER INFORMATION: motif
 90 <220> FEATURE:
 91 <221> NAME/KEY: VARIANT
 92 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15
 93 <223> OTHER INFORMATION: Xaa = Ala or Gly
 95 <400> SEQUENCE: 4
 96 Phe Xaa Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Thr
 97  1               5                   10                  15
100 <210> SEQ ID NO: 5
101 <211> LENGTH: 16
102 <212> TYPE: PRT
103 <213> ORGANISM: Artificial Sequence
105 <220> FEATURE:
106 <223> OTHER INFORMATION: motif
108 <220> FEATURE:
109 <221> NAME/KEY: VARIANT
110 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15
111 <223> OTHER INFORMATION: Xaa = Ala or Gly
113 <400> SEQUENCE: 5
114 Ser Xaa Xaa Xaa Xaa Asn Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Ile
115  1               5                   10                  15
118 <210> SEQ ID NO: 6
119 <211> LENGTH: 16
120 <212> TYPE: PRT
121 <213> ORGANISM: Artificial Sequence
123 <220> FEATURE:
124 <223> OTHER INFORMATION: motif
126 <220> FEATURE:
127 <221> NAME/KEY: VARIANT
128 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15
129 <223> OTHER INFORMATION: Xaa = Ala or Gly
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131 <220> FEATURE:
132 <221> NAME/KEY: VARIANT
133 <222> LOCATION: 16
134 <223> OTHER INFORMATION: Xaa = any amino acid
136 <400> SEQUENCE: 6
137 Ser Xaa Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Xaa
138 1   5   10   15

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 06/04/2002
PATENT APPLICATION: US/09/772,116   TIME: 12:41:12

Input Set : A:\Seqlist.txt
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16
Seq#:2; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15
Seq#:3; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15
Seq#:4; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15
Seq#:5; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15
Seq#:6; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15,16